

Untitled

RESULT 5

US- 10- 305- 278- 395

: Sequence 395, Application US/ 10305278

: Patent No. 7129338

GENERAL INFORMATION:

: APPLICANT: OTA, TOSHI O

: APPLICANT: ISOGAI, TAKAO

: APPLICANT: NISHIKAWA, TETSUO

: APPLICANT: KAWAI, YURI

: APPLICANT: SUGIYAMA, TOMOYASU

: APPLICANT: HAYASHI, KOJI

: TITLE OF INVENTION: SECRETORY PROTEIN OR MEMBRANE PROTEIN

: FILE REFERENCE: 084335/ 0121

: CURRENT APPLICATION NUMBER: US/ 10/ 305, 278

: CURRENT FILING DATE: 2002- 11- 27

: PRIOR APPLICATION NUMBER: US/ 09/ 611, 523

: PRIOR FILING DATE: 2000- 07- 07

: PRIOR APPLICATION NUMBER: JP 1999- 194179

: PRIOR FILING DATE: 1999- 07- 08

: PRIOR APPLICATION NUMBER: JP 2000- 118775

: PRIOR FILING DATE: 2000- 01- 11

: PRIOR APPLICATION NUMBER: JP 2000- 183766

: PRIOR FILING DATE: 2000- 05- 02

: PRIOR APPLICATION NUMBER: 60/ 159, 586

: PRIOR FILING DATE: 1999- 10- 18

: PRIOR APPLICATION NUMBER: 60/ 183, 323

: PRIOR FILING DATE: 2000- 02- 17

: NUMBER OF SEQ ID NOS: 679

: SOFTWARE: Patent In Ver. 2. 1

: SEQ ID NO 395

: LENGTH: 550

: TYPE: DNA

: ORGANISM Homo sapiens

: FEATURE:

: OTHER INFORMATION: "n" may be a, t, c, g, other or unknown

US- 10- 305- 278- 395

Query Match 38. 4% Score 483; DB 5; Length 550;

Best Local Similarity 97. 2% Pred. No. 2. 2e- 84;

Matches 486; Conservative 0; Mismatches 14; Indels 0; Gaps 0;

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Qy      1 ATGCTCCCCTGGA CGGCGCT CGGCGT GGCCCT GAGCTT GCGGCT GGCGCT GGCGCGGAGC 60
Db      51 ATGCTCCCCTGGA CGGCGCT CGGCGT GGCCCT GAGCTT GCGGCT GGCGCT GGCGCGGAGC 110
Qy      61 GGCGCGGAGCGCGGT CCA CCAGCAT CAGCCCCCGAGGGGACCT GAT GTT CCT GCT GGAC 120
Db     111 GGCGCGGAGCGCGGT CCA CCAGCAT CGGCCCCCGAGGGGACCT GAT GTT CCT GCT GGAC 170
Qy     121 AGCTCAGCCAGCGT CTCTCACTACGAGTTCTCCCGGGTT CGGGAGTTTGT GGGGCAGCTG 180
Db     171 AGCTCAGCCAGCGT CTCTCACTACGANTTCTCCCGGGTT CGGGAGTTTGT GGGGCAACTG 230
Qy     181 GTGGCTCACTGCCCCTGGGCA CCGGGGCGCT GCGT GCGAGT CTGGT GCACT GGGCAGT 240
Db     231 GTGGCTCACTGCCCCTGGGCA CCGGGGCGCT GCGT GCGAGT CTGGT GCACT GGGCAGT 290
Qy     241 OGGCCATACACGAGTTCCCTTT CGGCCAGCACAGCT OGGGT GAGGCT GCGCAGGAT GCG 300
Db     291 OGGCCATACACGAGTTCCCTTT CGGCCAGCACANCT OGGGT GAGGCT GCGCAGGAT GCG 350
Qy     301 GTGCGTGCTTCTGCCAGCGCATGGGTGACACCCACACTGGCTGGCGCTGGTCTATGCC 360
  
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U n t i t l e d

Db	351	GTGCGTGCTTCTGCCANOGCATGGGTGACACCCACACTGGCCTGGGCGCTGGTCTATGOC	410
Qy	361	AAGGAACAGCTGTTTGCTGAAGCATCAGGTGCGCGGCCAGGGGTGCCAAAAGTGCTGGTG	420
Db	411	AAGGAACAGCTGTTTGCTGAANCATCAGGTGCGCGGCCAGGGGTGCCAAAAGTGCTGGTG	470
Qy	421	TGGGTGACAGATGGGGGCTCCAGCGACCCCTGTGGGCCCCCCCCATGCAGGAGCTCAAGGAC	480
Db	471	TGGGTGACANATGGGGGCTCCNCGACCCCTGTGGGCCCCCCCCATGCACGANCTCAAGGAC	530
Qy	481	CTGGGCGTCACCGTGTTTCAT	500
Db	531	CTGGGCNTCACCGTGTCAT	550